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RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/008,277A
Source: OIPE
Date Processed by STIC: 6-3-02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>>, EFS Submission User Manual - ePAVE).
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 10/008, 277A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 1 Wrapped Amino
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 3 Numbering
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0 A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 6 "bug"
- 7 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(OLD RULES) (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
- 7 (OLD RULES)
- 7 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
(NEW RULES) <210> sequence id number
<400> sequence id number
000
- 8 (NEW RULES)
- 9 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
(NEW RULES) Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213> Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or
Response scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0 Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
"bug" resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

Does Not Comply
Corrected Diskette Needed



OIPE

RAW SEQUENCE LISTING

DATE: 06/03/2002

PATENT APPLICATION: US/10/008,277A

TIME: 13:24:42

Input Set : A:\VPI99-06 CON.ST25.txt

Output Set : N:\CRF3\06032002\J008277A.raw

3 <110> APPLICANT: Vertex Pharmaceuticals Incorporated
 4 Salituro, Francesco
 5 Bemis, Guy
 6 Green, Jeremy
 7 Jasna, Fejzo
 8 Xie, Xiaoling
 10 <120> TITLE OF INVENTION: Inhibitors of c-Jun N-Terminal Kinases (JNK)
 12 <130> FILE REFERENCE: VPI/99-06 CON
 14 <140> CURRENT APPLICATION NUMBER: 10/008,277A
 C--> 15 <141> CURRENT FILING DATE: 2002-05-17
 17 <150> PRIOR APPLICATION NUMBER: PCT/US00/15248
 18 <151> PRIOR FILING DATE: 2000-06-02
 20 <150> PRIOR APPLICATION NUMBER: 60/237,523
 21 <151> PRIOR FILING DATE: 1999-06-02
 23 <160> NUMBER OF SEQ ID NOS: 3
 25 <170> SOFTWARE: PatentIn version 3.0
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 39
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Other
 32 <400> SEQUENCE: 1
 34 Gly Cys Thr Cys Thr Ala Gly Ala Gly Cys Thr Cys Cys Ala Thr Gly
 35 1 5 10 15
 37 Gly Gly Cys Ala Gly Cys Ala Ala Ala Gly Cys Ala Ala Gly
 38 20 25 30
 40 Thr Thr Gly Ala Cys Ala Ala
 41 35
 43 <210> SEQ ID NO: 2
 44 <211> LENGTH: 37
 45 <212> TYPE: PRT
 46 <213> ORGANISM: Other
 48 <400> SEQUENCE: 2
 50 Thr Ala Gly Cys Gly Gly Ala Thr Cys Cys Thr Cys Ala Thr Thr Cys
 51 1 5 10 15
 53 Thr Gly Ala Ala Thr Thr Cys Ala Thr Thr Ala Cys Thr Thr Cys Cys
 54 20 25 30
 56 Thr Thr Gly Thr Ala
 57 35
 59 <210> SEQ ID NO: 3
 60 <211> LENGTH: 21
 61 <212> TYPE: PRT
 62 <213> ORGANISM: Other
 64 <400> SEQUENCE: 3

See item #10 on
ERROR Summary SHEET

RAW SEQUENCE LISTING

DATE: 06/03/2002

PATENT APPLICATION: US/10/008,277A

TIME: 13:24:42

Input Set : A:\VPI99-06 CON.ST25.txt

Output Set: N:\CRF3\06032002\J008277A.raw

66 Lys Arg Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn
67 1 5 10 15
69 Gln Ala Leu Leu Arg
70 20

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/008,277A

DATE: 06/03/2002

TIME: 13:24:43

Input Set : A:\VPI99-06 CON.ST25.txt

Output Set: N:\CRF3\06032002\J008277A.raw

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date